# Special Session on Soft Computing in Computational Biology & Bioinformatics (SoCCoBB 2009)

In conjunction with the International Conference on Soft Computing and Pattern Recognition (SoCPaR 2009), Melaka December 4 – 7, 2009

#### **Session Organizers:**

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# **Call For Paper**

Bioinformatics and computational biology present a number of difficult problems with large search spaces. Recent applications of Soft Computing methods in this area suggest that they are well-suited to this area of research. This special session will highlight applications of Soft Computing to a broad range of topics. Particular interest will be directed towards novel applications of Soft Computing to problems in these areas. The following areas are the topics of interests but not limited to:

- Knowledge Discovery and Data Mining techniques for the Analysis of high throughput biological data
- Mathematical modeling of Evolutionary Computing
- Bayesian approaches, Neural Networks, Support Vector Machine, Case-Based Reasoning
- o Microarray design and analysis
- o Visualization of biological systems and networks
- Data and software integration
- o Sequence analysis
- Gene Expression and gene regulatory networks
- Phylogenetics and phylogenomics
- Comparison and Alignment methods
- Feature Selection

# SUBMISSION GUIDELINES

Papers are invited from prospective authors with interests in this particular special session and related areas of applications. All contributions should be original and not published elsewhere or intended to be published during the review period. Details of the required paper format are published at the official SoCPaR 2009 website: (http://www.socpar.org/)

# **Important Dates:**

Deadline for paper submission: June, 26, 2009 Notification of acceptance/rejection: August 14, 2009 Deadline for camera-ready manuscript submission: September 14, 2009